

# Emerging roles of growth differentiation factor-15 in brain disorders (Review)

WEI-WEI JIANG<sup>1\*</sup>, ZI-ZHEN ZHANG<sup>2\*</sup>, PING-PING HE<sup>3,4\*</sup>, LI-PING JIANG<sup>1,5</sup>, JIN-ZHI CHEN<sup>1</sup>,  
XING-TING ZHANG<sup>1</sup>, MI HU<sup>1</sup>, YANG-KAI ZHANG<sup>1</sup> and XIN-PING OUYANG<sup>1,4\*</sup>

<sup>1</sup>Department of Physiology, Institute of Neuroscience Research, Hengyang Key Laboratory of Neurodegeneration and Cognitive Impairment, Hengyang Medical College, University of South China; <sup>2</sup>Department of Medical Humanities, School of Medicine, Hunan Polytechnic of Environment and Biology; <sup>3</sup>Hunan Province Cooperative Innovation Centre for Molecular Target New Drug Study, Nursing School, University of South China; <sup>4</sup>Institute of Cardiovascular Research, Key Laboratory for Atherosclerosis of Hunan Province, Hunan Province Cooperative Innovation Center for Molecular Target New Drug Study, University of South China, Hengyang, Hunan 421001; <sup>5</sup>Department of Critical Care Medicine, Hunan Taihe Hospital, Changsha, Hunan 410004, P.R. China

Received March 5, 2021; Accepted August 6, 2021

DOI: 10.3892/etm.2021.10705

**Abstract.** Brain disorders, such as Alzheimer's and Parkinson's disease and cerebral stroke, are an important contributor to mortality and disability worldwide, where their pathogenesis is currently a topic of intense research. The mechanisms underlying the development of brain disorders are complex and vary widely, including aberrant protein aggregation, ischemic cell necrosis and neuronal dysfunction. Previous studies have found that the expression and function of growth differentiation factor-15 (GDF15) is closely associated with the incidence of brain disorders. GDF15 is a member of the TGF $\beta$  superfamily, which is a dimer-structured stress-response

protein. The expression of GDF15 is regulated by a number of proteins upstream, including p53, early growth response-1, non-coding RNAs and hormones. In particular, GDF15 has been reported to serve an important role in regulating angiogenesis, apoptosis, lipid metabolism and inflammation. For example, GDF15 can promote angiogenesis by promoting the proliferation of human umbilical vein endothelial cells, apoptosis of prostate cancer cells and fat metabolism in fasted mice, and GDF15 can decrease the inflammatory response of lipopolysaccharide-treated mice. The present article reviews the structure and biosynthesis of GDF15, in addition to the possible roles of GDF15 in Alzheimer's disease, cerebral stroke and Parkinson's disease. The purpose of the present review is to summarize the mechanism underlying the role of GDF15 in various brain disorders, which hopes to provide evidence and guide the prevention and treatment of these debilitating conditions.

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*Correspondence to:* Dr Xin-Ping Ouyang, Department of Physiology, Institute of Neuroscience Research, Hengyang Key Laboratory of Neurodegeneration and Cognitive Impairment, Hengyang Medical College, University of South China, 28 Changsheng West Road, Hengyang, Hunan 421001, P.R. China  
E-mail: yl655@163.com

\*Contributed equally

**Abbreviations:** GDF15, growth differentiation factor-15; GFRAL, GDNF receptor  $\alpha$ -like; AD, Alzheimer's disease; PD, Parkinson's disease; EGR-1, early growth response-1; lncRNAs, long non-coding RNAs; OSCC, oral squamous cell carcinoma; HCC, hepatocellular carcinoma; miRNAs, microRNAs; CCN2, connective tissue growth factor; MSA, methylseleninic acid; ARRBI,  $\beta$ -arrestin1; LPS, lipopolysaccharide; HRV, human rhinovirus; TGF $\beta$ RII, TGF $\beta$  receptor type II; A $\beta$ , amyloid  $\beta$ ; 6-OHDA, neurotoxin 6-hydroxydopamine; hUCB-MSCs, human umbilical cord blood-derived mesenchymal stem cells

**Key words:** growth differentiation factor-15, brain disorders, Alzheimer's disease, cerebral stroke, Parkinson's disease

## Contents

1. Introduction
2. Structure and biosynthesis of GDF15
3. Regulation of GDF15 expression
4. Biological functions of GDF15
5. Role of GDF15 in major brain disorders
6. Conclusion and future directions

## 1. Introduction

Brain disorders, along with malignant tumors and cardiovascular disorders, are among the leading causes of morbidity and mortality worldwide (1). Brain disorders pose a serious socioeconomic burden, where the Global Burden of Disease 2017 data demonstrated that ~324.4 million individuals were affected by brain disorders in Europe, which accounts for

79.1% of all non-communicable diseases in 2017 (1). Although cholinesterase inhibitors and dopamine-like drugs have shown considerable efficacy for the treatment of brain disorders, particularly in patients with degenerative diseases in the central nervous system, side effects and long-term sequelae remain (2). Therefore, a more detailed understanding in the mechanism underlying the progression and pathophysiology of brain disorders is crucial for developing new therapeutic strategies.

Growth differentiation factor-15 (GDF15), which was first identified in the human cDNA library that was enriched for genes associated with macrophage activation using the subtraction cloning approach, is a distant member of the TGF $\beta$  superfamily (3). GDF15 is highly expressed in the heart, liver, kidney, intestine, lung, placenta and the prostate gland (4-6). In humans, the physiological concentration of GDF15 lies in the range of 200-1,200 pg/ml, where its levels increase with age (7). It has been reported that GDF15 participates in tissue repair by exerting antiapoptotic and anti-inflammatory effects whilst maintaining vascular endothelial functions (8,9). There is also accumulating evidence that GDF15 is involved in the occurrence and development of cardiovascular diseases, diabetes and cancer (10-12). In 2017, research successively identified glial cell-derived neurotrophic factor receptor  $\alpha$ -like (GFRAL) as the receptor of GDF15 (13-16). Discovery of the GDF15/GFRAL signaling pathway provided a potentially novel target for the treatment of obesity and metabolic diseases (16-18). In recent years, accumulating evidence has also indicated that GDF15 is associated with a range of brain disorders, including Alzheimer's disease (AD), cerebral stroke and Parkinson's disease (PD) (19-21). Therefore, the aim of the present review is to summarize the regulatory processes and physiological functions of GDF15, with an emphasis on its role in brain disorders.

## 2. Structure and biosynthesis of GDF15

The GDF15 gene can be mapped onto the chromosome 19p13.1-13.2 genomic locus, which contains two exons and one single intron with an open reading frame of 924 bp (3). Its corresponding mRNA can be translated into a 308-amino acid polypeptide, which is composed of the following three parts: Signal peptide, pro-peptide region and a mature region on the carboxyl terminus (3). The mature domain of 112 amino acids is first separated from the propeptide region of 167 amino acids by a furin-like convertase, which is then cleaved by furin/paired basic amino-acid-cleaving enzyme 4 and MMP-26 (22,23). The mature domain of GDF15 contains a highly conserved pattern of seven cysteine residues, where six of these form intra-chain disulfide bonds, forming a highly stable cysteine structure that is resistant to physical and chemical damage, including enzymatic attacks (24). Unlike other TGF $\beta$  families of proteins, the propeptide is not required for proper GDF15 folding (25). Although propeptides may facilitate the detection of improper GDF15 folding, engineered GDF15 that lacks the pro-peptide domain can still be secreted in its proper folded form (26). GDF15 is generally secreted as a dimer that is formed by inter-chain disulphide bonds, which performs complex biological functions through autocrine or paracrine pathways (Fig. 1) (25). In addition, GDF15 can be

secreted as an unprocessed pro-peptide, where it can bind to the extracellular matrix (ECM) through its 89 amino acids on the C-terminal domain in a reversible manner in prostate cancer (27). There, GDF15 can be released from the ECM into the circulation by locally acting MMPs or pro-convertases (27).

## 3. Regulation of GDF15 expression

GDF15 is a type of stress-response protein (28). It was previously reported that GDF15 expression is markedly increased under conditions of inflammation, ischemia, hypoxia and organ damage (29). GDF15 is important for the regulation of angiogenesis, apoptosis, lipid metabolism and inflammation, some of the factors that have been found to regulate GDF15 expression are discussed in this section.

*p53*. As a tumor suppressor gene, *p53* serves a key role in controlling cell proliferation, inhibiting malignant cell proliferation and regulating cell cycle progression (30). *p53* was one of the first transcription factors that was identified to be transcriptional regulators of GDF15 expression (31). There are  $\geq$  two *p53* binding sites in the GDF15 promoter, one near the transcription start site and another in the region that lie 851 bp upstream of the transcription start site (31), where both binding sites can transactivate the GDF15 promoter (31). It was previously reported that GDF15 expression was robustly upregulated following the overexpression or pharmacological induction of *p53* in human lung cancer cell lines, osteosarcoma cell lines and breast cancer cell lines (32,33). In addition, the DNA intercalator doxorubicin was found to significantly increase GDF15 expression in *p53* wild-type human breast cancer cell lines, but exerted no effects on *p53*-null cells (34). *In vitro*, GDF15 expression was found to be markedly increased in human bronchial epithelial cells and human pulmonary vascular endothelial cells upon exposure to hyperoxia, whilst *p53* knockdown robustly reduced this induction of GDF15 transcription by hyperoxia (35). Therefore, this suggests that *p53* is an important regulator of the production of GDF15. Similarly, C-reactive proteins in human aortic endothelial cells and human maternal expression gene 3 in colon cancer cell lines were both shown to induce GDF15 expression by recruiting the *p53* protein (36,37). Different members of the *p53* tumor suppressor gene family have been demonstrated to exhibit different binding affinities for GDF15 (38). The GDF15 promoter region contains two *p53*-type response elements (RE), RE1 and RE2, where most 3' quarter-sites (areas bound by the *p53* tetramer) in RE2 exhibit a higher binding affinity for *p53* (38). Therefore, GDF15 is activated by *p53* to a greater extent compared with that by other related family members, including *p63* and *p73* (38).

*Early growth response-1 (EGR-1)*. *EGR-1* is a transcription factor that contains three zinc finger domains and is considered to be an important regulator of GDF15 (39). A number of studies have previously shown that pharmacological agonists, such as troglitazone and non-steroidal anti-inflammatory drugs, can increase *EGR-1* expression in human colon cancer cell lines prior to GDF15 induction, supporting the hypothesis that GDF15 is a downstream target of *EGR-1* (40-42). Indeed, position -73 to -51 of the GDF15 promoter contains

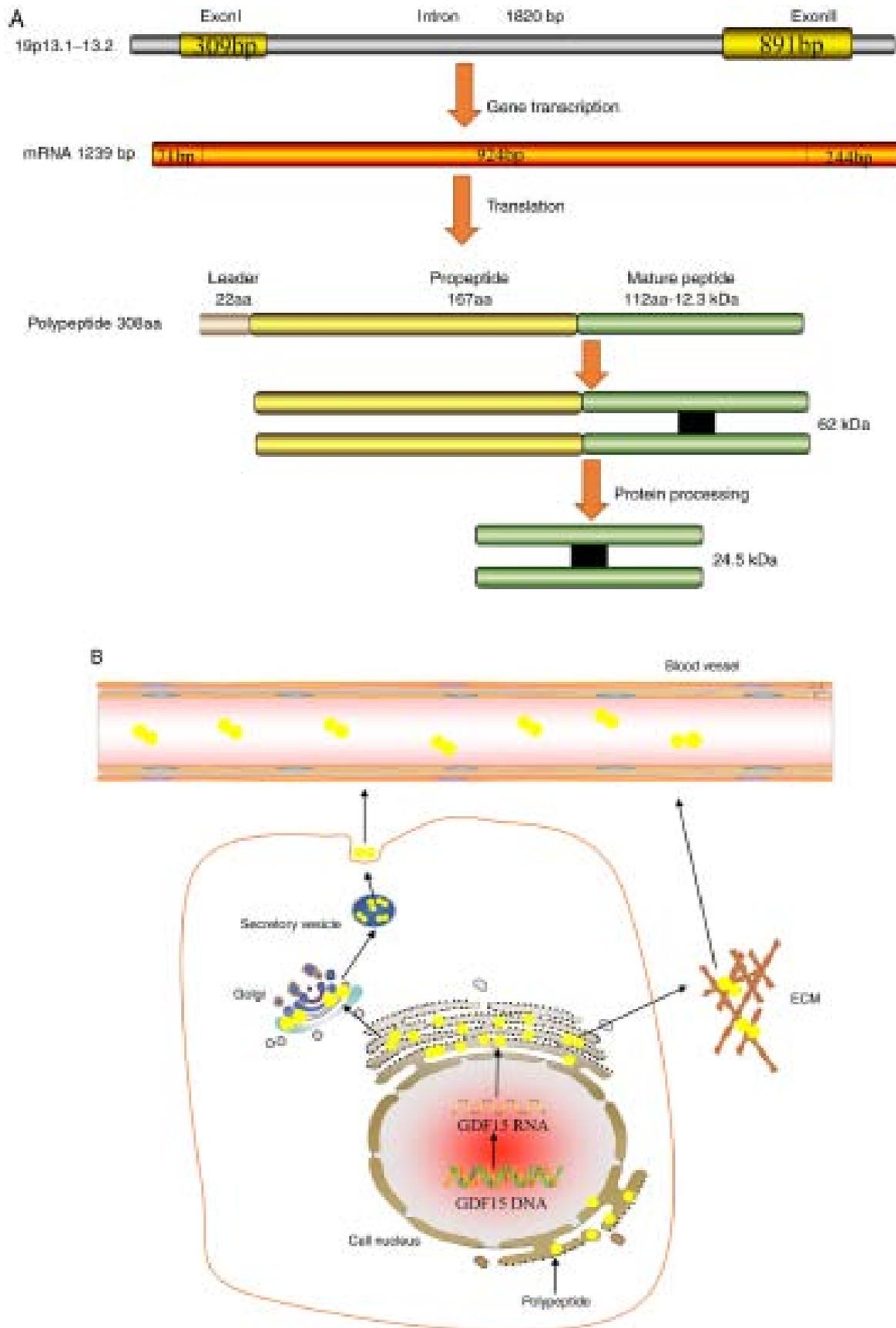


Figure 1. Synthesis and Secretion of GDF15. (A) GDF15 gene that is located on the chromosome 19p13.1-13.2 genomic locus can produce a mRNA of 1239 BP. And the mRNA is translated into a 308 amino acid polypeptide, which includes signal peptide, pro-peptide region and a mature region. Then the mature domain is separated from the propeptide region by a furin-like convertase. (B) After transcription, translation and processing, GDF15 (yellow circle) is formed into a 62-kDa protein, which is transported into the circulation through vesicles. In addition, the unprocessed GDF15 can also be directly secreted to bind to ECM. After hydrolysis by metalloenzymes or pro-convertases, GDF15 enters the circulation from ECM. GDF15, growth differentiation factor-15; ECM, extracellular matrix.

EGR-1-binding sites (43). Furthermore, DNA methylation at the -53 site of the GDF15 promoter site blocks the binding of EGR-1, which subsequently inhibits GDF15 induction *in vitro* (43). By contrast, GDF15 transcription by EGR-1 can be restored following incubation with 5-aza-2'-deoxycytidine (a demethylating reagent) (43). Accordingly, in HT29 colon carcinoma cells, activity of the GDF15 promoter and GDF15 expression are markedly increased by the ectopic expression of EGR-1 in a dose-dependent manner, whereas EGR-1 knockdown using small interfering (si)RNAs was shown to significantly decrease silibinin-induced GDF15 expression (44). These observations suggest that EGR-1 is a direct transcriptional regulator of GDF15.

*Long non-coding RNAs (lncRNAs) and microRNAs (miRNAs/miRs).* LncRNAs represents a class of RNAs that cannot be translated into proteins and are typically >200 nucleotides in length (45). It has been reported extensively that lncRNAs can serve key roles in the regulation of gene expression. Kong *et al* (46) previously revealed that LINC0113 overexpression decreased the mRNA and protein expression levels of GDF15 in oral squamous cell carcinoma (OSCC) cell lines, whilst treatment with the exogenous recombinant human GDF15 protein was able to restore the migratory and invasive abilities of OSCC cells that was previously weakened by LINC0113. In addition, a significant positive correlation was identified between lncRNA plasmacytoma variant translocation 1 (PVT1) expression and GDF15 expression in hepatocellular carcinoma (HCC) tissues (47). GDF15 knockdown using siRNAs significantly suppressed the proliferation of HCC cells caused by lncRNA PVT1 overexpression, suggesting that lncRNA PVT1 may be an important upstream regulator of GDF15 expression in HCC cells (47). Another study also revealed that CCAAT/enhancer-binding protein  $\beta$  (CEBPB) can bind to the promoter of GDF15 to facilitate GDF15 gene expression in ovarian cancer cell lines (48). In addition, GDF15 expression was previously found to be negatively associated with that of the lncRNA growth arrest-specific 5 (GAS5) in ovarian cancer tissues (48). Mechanistically, lncRNA GAS5 was shown to competitively bind to CEBPB to inhibit the promoting effect of CEBPB on GDF15 transcription (48). It is expected that additional lncRNAs involved in the regulation of GDF15 expression will be discovered by future studies.

Similar to lncRNAs, miRNAs are also an important component of gene transcription regulators, which functions by pairing with the 3'-untranslated region of target mRNAs (49). Both miR-873-5p and miR-1233-3p have been shown to exert suppressive effects on GDF15 expression in melanoma cell lines (50). However, single-nucleotide polymorphism in miRNA, rs1054564, was located in the GDF15'UTR complementary to the hsa-miR-1233-3p seed region, and the presence of this C-allele was discovered to weaken the binding of hsa-miR-1233-3p to GDF15, thereby enhancing the expression of the GDF15 protein (50). miR-3189 is a primate-specific miRNA that is embedded within the introns of GDF15 (51). Increased miR-3189 expression results in elevated GDF15 expression by down-regulating p53 in colorectal cancer cell lines (51). In addition, overexpression of miR-3189 in HCT116-p53<sup>-/-</sup> colorectal cancer cells was shown to upregulate the expression of a

subset of p53 targets, including GDF15 and growth arrest and DNA-damage-inducible 45 $\alpha$  (51).

*Hormones and hormone derivatives.* Hormones and hormone derivatives have also been demonstrated to lie upstream of GDF15 (52). Primary adrenal insufficiency is accompanied by an increase in GDF15 expression, where glucocorticoid replacement therapy was shown to effectively reduce this concentration of GDF15 in a dose-dependent manner (53). In brown adipose tissues, noradrenergic cAMP-mediated thermogenic activation was found to increase GDF15 gene expression and subsequent release (54). In addition, metformin was reported to increase GDF15 levels, but it had no effect on GFRAL receptor-deficient mice (52). Zhao *et al* (55) previously found that thyroid hormone levels were independently associated with GDF15 expression, such that T3 treatment promoted GDF15 expression in brown adipose tissues of C57BL/6 mice. Furthermore, previous *in vivo* and *in vitro* experiments demonstrated that testosterone and estradiol treatment reduced GDF15 secretion through androgen receptor/estrogen receptor-mediated signaling pathways (56).

#### 4. Biological functions of GDF15

*Angiogenesis.* Accumulating evidence has suggested that GDF15 is a potential stimulator of angiogenesis (Table I). After being secreted by senescent endothelial colony-forming cells generated from adult human blood in a paracrine manner, GDF15 can promote proliferation, migration and NO production in non-senescent endothelial colony-forming cells generated from adult human blood (57). During this process, a number of signaling pathways are activated by GDF15 in an oxidative stress-independent manner, including AKT, ERK1/2 and Smad2 (57). This improved the function of vascular progenitor cells, which may serve therapeutic effects on the damaged vascular system (57). Similarly, GDF15 can also enhance the expression of cyclins D1 and E in HUVECs through the PI3K/AKT, JNK and ERK signaling pathways to promote the proliferation of endothelial cells (58). GDF15 is also involved in the mechanism underlying ischemia/reperfusion injury. During the process of cardiac ischemia, GDF15 was shown to stimulate the angiogenesis of hypoxic HUVECs by inhibiting p53 signaling whilst upregulating hypoxia-inducible factor 1 $\alpha$  expression (59). Furthermore, since the repair of large bone defects remains to be a major medical challenge, GDF15 was shown to represent a potentially effective solution. Wang *et al* (60) found that GDF15 can promote the formation of functional blood vessels at the site of artificially-induced angiogenesis, which significantly improved the healing of critical-sized calvarial defects. Neovascularization is one of the major characteristics in cancer (61). Following chemotherapy, the expression of GDF15 was found to be markedly upregulated in HCC (62). GDF15 can induce Src and then activate AKT, MAPK and NF- $\kappa$ B downstream in HCC, which promotes the proliferation, migration and tube formation of surrounding endothelial cells *in vitro* (62). By contrast, thalidomide, an agent with known anti-angiogenic activities, can significantly attenuate and reverse these effects aforementioned (62). In addition, it was found that GDF15 interacted with connective tissue growth factor (CCN)-2, to inhibit

Table I. Biological functions of GDF15.

Actions	Experimental models	Mechanisms	(Refs.)
<b>Angiogenesis</b>			
Pro-angiogenesis	Endothelial colony forming cells generated from adult human blood	↑NO, ↑AKT, ERK1/2 and SMAD2	(57)
	HUVEC	↑Cyclin D1 and E, ↑retinoblastoma phosphorylation and E2F-1 nuclear translocation	(58)
	HUVEC	↓p53, hypoxia-induced factor-1 $\alpha$	(59)
	Hepatocellular carcinoma	↑Src and AKT, MAPK and NF- $\kappa$ B downstream	(62)
Anti-angiogenesis	HUVEC	↓Connective tissue growth factor 2/focal adhesion kinase	(63)
<b>Apoptosis</b>			
Pro-apoptosis	Prostate cancer cells	↑Methylseleninic acid, ↑caspase-dependent apoptosis	(66,67)
	Activated macrophages	↑ PARP, caspase-3 or AIF	(68)
	A459 cells	↑caspase-9 and caspase-3; ↓ERK1/2 and p38 MAPK phosphorylation	(69)
Anti-apoptosis	HUVEC	↑PI3K/AKT/ endothelial nitric oxide synthase; ↓NF- $\kappa$ B/JNK	(70)
	Patients with pulmonary hypertension	↑AKT	(71)
<b>Lipid metabolism</b>			
Lipid metabolism	Transgenic GDF15 mice	↑Key thermogenic and lipolytic genes	(73)
	Mice fed with high-fat diet	↑Glial cell-derived neurotrophic factor receptor $\alpha$ -like	(74)
	Patients with non-alcoholic steatohepatitis	↑ $\beta$ -arrestin1; ↑ $\beta$ -oxidation genes; ↓fatty acids	(75)
	Fasted mice	↑Fatty acid $\beta$ -oxidation and ketogenesis	(76)
<b>Inflammation</b>			
Anti-inflammation	Mice treated with lipopolysaccharide	↓Monocyte chemoattractant protein-1, TNF- $\alpha$ , transforming growth factor- $\beta$ -activated kinase 1 phosphorylation and NF- $\kappa$ B	(79,81)
	Mice	↑Triglyceride metabolism	(77)
Pro-inflammation	Human GDF15 transgenic mice	↓IFN- $\lambda$ 2/3 mRNA	(78)
	Human transgenic mice tracheal epithelial cells	↓IFN- $\lambda$ 1/IL-29	(78)

↑, enhancement or promotion; ↓, reduction or inhibition; GDF-15, growth differentiation factor-15; HUVECs, human umbilical cord vascular endothelial cells; IFN, interferon.

CCN2-mediated focal adhesion kinase activation, which in turn decreased  $\alpha$ v $\beta$ 3 integrin clustering in HUVECs, to exert antagonistic effects on angiogenesis (63). This phenomenon is conducive to understanding the role of GDF15 under various disease conditions further.

**Cell apoptosis.** Apoptosis is a process in which cell death occurs in an orderly manner and is crucial for the maintenance of internal homeostasis (64). Owing to its reported function as a stress-response protein, GDF15 has been reported to regulate apoptosis (65). GDF15 is a downstream target of methylseleninic acid (MSA), where GDF15 knock-down significantly inhibited the apoptosis of prostate cancer cells mediated by MSA (66,67). By contrast, GDF15 overexpression made prostate cancer cells flattened and more spread out and induced caspase-dependent apoptosis (66,67). GDF-15 was inducible in human macrophages by oxidized low density lipoprotein and its mediators *in vitro*, and GDF15

immunoreactivity was colocalized with apoptosis markers such as PARP, caspase-3 or apoptosis-inducing factor immunoreactivity, suggesting that GDF15 may modulate apoptosis process in activated macrophages (68). In addition, A549 lung adenocarcinoma cell apoptosis was also induced by GDF15 overexpression through promoting caspase-9 and caspase-3 expression and inhibition of ERK1/2 and p38 activation, which was mediated in a TGF $\beta$  receptor type II (TGF $\beta$ RII)-dependent manner (69). Conversely, GDF15 can exert a protective effect against the apoptosis of HUVECs induced by high glucose concentrations (70). Mechanistically, this effect may be caused by GDF15 maintaining the activity of PI3K/Akt/eNOS pathway and attenuating NF- $\kappa$ B/JNK pathway (70). In addition, under conditions of hypoxia and laminar shear stress, GDF15 expression in the pulmonary microvascular endothelial cells of patients with pulmonary arterial hypertension was found to be significantly higher compared with that in normal lung tissues, where the extent

of cell apoptosis was reduced by GDF15 overexpression in an AKT-dependent manner (71).

**Lipid metabolism.** GDF15 can serve as a potential lipid metabolism regulator (72). GDF15 overexpression conferred higher resistance to diet- and genetic-induced obesity in transgenic GDF15 mice compared with that in wild-type mice by increasing lipid oxidation whilst promoting the expression of thermogenic genes and adipose tissue metabolism (73). Following treatment with the GDF15 antibody, the weight, obesity and the degree of liver lipid deposition of mice fed on a high-fat diet were markedly higher compared with those in the control group (74). Therefore, GDF15 may be a potential target for the treatment of fatty liver disease. Zhang *et al* (75) revealed that  $\beta$ -arrestin 1 (ARRB1) deficiency in patients with non-alcoholic steatohepatitis was accompanied with increased free fatty acid levels and decreased  $\beta$ -oxidation gene expression in a GDF15-dependent manner in liver. In addition, ARRB1 was found to interact with GDF15 to promote the translocation of the GDF15 precursor to the Golgi body for cleavage and maturation (75). During fasting, the inositol-requiring enzyme 1 $\alpha$ /X-box binding protein axis can increase the expression of GDF15 by binding to its promoter to promote fatty acid oxidation and ketone production in the liver (76). However, GDF15 knockout can significantly suppress  $\beta$ -oxidation and ketogenesis in mice with streptozocin-induced type I diabetes or in mice subjected to fasting (76).

**Inflammatory response.** In addition to the aforementioned functions, GDF15 can also exert anti-inflammatory and proinflammatory properties (77,78). GDF15 can inhibit the inflammatory response induced by lipopolysaccharide (LPS) (79). A previous study demonstrated that GDF15-knockout mice displayed worsened characteristics following the induction of LPS-induced renal and cardiac injury, whilst GDF15 overexpression conferred opposite effects (80). GDF15 can inhibit the activation of the NF- $\kappa$ B pathway to reduce the production of proinflammatory factors, including monocyte chemoattractant protein-1 and TNF- $\alpha$  (81). This in effect prevents LPS-induced liver injury in mice by blocking the phosphorylation of TGF $\beta$ -activated kinase 1 (81). Luan *et al* (77) previously reported that GDF15 improved cardiac and hepatic tolerance to inflammation in mice by regulating triglyceride metabolism. GDF15 was also found to regulate the response to human rhinovirus (HRV) infection and virus-induced lung inflammation (78). In human GDF15 transgenic mice, the overexpression of GDF15 resulted in enhanced inflammatory responses to HRV and decreased IFN- $\lambda$ 2/3 mRNA expression (78). In addition, the IFN- $\lambda$ 1/IL-29 protein, which has antiviral activity, was found to be inhibited by GDF15 in tracheal epithelial cells from human GDF15 transgenic mice, which promoted HRV replication and the subsequent inflammatory response (78).

## 5. Role of GDF15 in major brain disorders

**AD.** AD is a progressive neurodegenerative disease that mainly manifests with clinical symptoms of cognitive and behavioral impairment (82). The pathogenesis of AD is complex and may involve amyloid  $\beta$  (A $\beta$ ) accumulation, tau

protein toxicity, synaptic damage, mitochondrial dysfunction and oxidative stress (83). GDF15 expression can be detected in the adult rat central and peripheral nervous systems, where it is mainly secreted into the cerebrospinal fluid (84,85). It has been demonstrated that GDF15 is closely associated with cognitive impairment, which is a major characteristic of AD (86). It was also previously shown that cognitive impairments due to dementia and AD were associated with higher GDF15 levels, particularly in the presence of cerebrovascular disease (87). By contrast, GDF15-deficient mice were shown to exhibit superior fear-associated memory and sensorimotor gating, which is conducive to cognition (88). Decreased numbers of white matter and grey matter nerve fibers, coupled with the increased volume of atrophy, may be important pathophysiological features of AD (89). Jiang *et al* (90,91) revealed that higher GDF15 levels were negatively correlated with gray matter volume and white matter integrity. In addition, GDF15 expression likely exhibits a close association with learning and memory impairments, where the hippocampus is the region that is typically worst affected by AD (86,92). GDF15-knockout mice were shown to display a progressive loss of motor neurons coupled with the decreased proliferation and migration of adult hippocampal progenitor cells (93,94). This appears to be a result of the absence of epidermal growth factor receptor signaling stimulation, which is normally promoted by GDF15 in a CXC chemokine receptor 4-dependent manner (93,94). Kim *et al* (95) reported that human recombinant GDF15 can enhance the proliferation and synaptic activity of mouse hippocampal neural stem cells both *in vitro* and *in vivo*, whereas GDF15 knockdown can reduce the proliferation of hippocampal neural stem cells *in vitro*. Within the neural network, the synapse forms a key component in mediating signal transmission between neurons, which underlies the mechanism of the generation and retention of memories (96). It was previously reported that GDF15 promoted synaptic glutamate release and increased the miniature excitatory post-synaptic current frequency in the medial prefrontal cortex of mice (97). The potential mechanism was mediated through activation of TGF $\beta$ RII-mediated ERK1/2 signaling to promote Ca $_v$ 3.1 and Ca $_v$ 3.3  $\alpha$  subunit expression, which increases T-type calcium channel activity (97). This suggests that GDF15 deficiency may impact synaptic function and accelerate AD progression (97). The production and accumulation of A $\beta$  is one of the main mechanisms underlying AD development (98). TGF $\beta$ RII is mainly expressed in the microglia and neurons, where it participates in GDF15-associated signaling pathways including Akt/mTOR pathway. GDF15 as a soluble paracrine factor can act on microglial cells to increase the expression level of TGF $\beta$ RII during AD (12). In addition, GDF15 can enhance the activity of insulin-degrading enzyme which, together with TGF $\beta$ RII, can promote A $\beta$  protein clearance (12). Decreased expression levels of TGF $\beta$ RII were found in human and mouse models of AD, which may be the underlying cause of the increased A $\beta$  accumulation and neurodegeneration (Fig. 2) (99,100). In summary, GDF15 appears to be involved in AD not only by promoting hippocampal neurogenesis and synaptic activity, but also by enhancing the clearance of the A $\beta$  protein. Therefore, GDF15 may represent an attractive therapeutic target for AD.

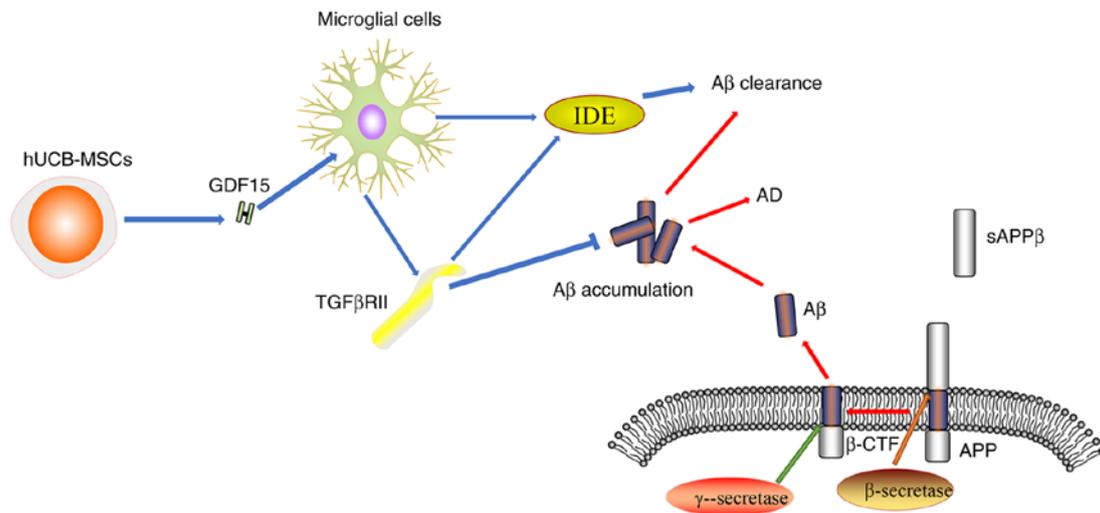


Figure 2. Association between GDF15 and AD. APP protein is hydrolyzed by  $\beta$ -secretase. SAPP $\beta$  is released into the extracellular matrix and  $\beta$ -CTF and A $\beta$  are produced by  $\gamma$ -Secretase. A $\beta$  accumulation is a leading cause of AD. hUCB-MSCs secrete GDF15, which can increase the expression of TGF $\beta$ RII on microglia through a paracrine mechanism, TGF $\beta$ RII in turn exerts a protective effect by inhibiting A $\beta$  accumulation. TGF $\beta$ RII can also increase the expression of IDE induced by GDF15. IDE is a degradation enzyme that can promote the clearance of A $\beta$ . GDF15, growth differentiation factor-15; AD, Alzheimer's disease; A $\beta$ , amyloid  $\beta$ ; hUCB-MSCs, human umbilical cord blood-derived mesenchymal stem cells; TGF $\beta$ RII, TGF $\beta$  receptor type II; IDE, insulin-degrading enzyme; SAPP $\beta$ , soluble peptide APP $\beta$ ; APP, amyloid precursor protein;  $\beta$ -CTF,  $\beta$ -secretase-derived fragment, C99.

**Cerebral stroke.** Cerebral stroke is an acute cerebrovascular disease, the pathogenesis of which is characterized by inadequate blood flow to the brain due to the rupture or occlusion of cerebrovascular vessels, thereby causing brain damage (101). Stroke can be divided into two categories, namely ischemic and hemorrhagic stroke, where American Stroke Association data in 2018 indicated that ischemic stroke accounted for 87% of all cases (102). Since GDF15 levels were found to be elevated after tissue injury, ischemia or hypoxia, it was hypothesized that there may be an association between GDF15 and stroke. Xiang *et al* (103) demonstrated that the genotype and allele frequencies of the GDF15 rs1804826G/T polymorphism were associated with the risk of cerebral stroke within the Chinese population. In a prospective, nested, case-controlled study of 27,628 initially healthy female individuals, Brown *et al* (104) revealed that the GDF15 concentration in patients with cardiovascular events, including stroke, was higher compared with that in individuals without cardiovascular events. In addition, a concentration higher than the 90th percentile (>856 pg/ml) was associated with a 2.7-fold increase in the risk of developing cardiovascular events (104). Previous studies reported that GDF15 levels may serve as a prognostic marker in patients with a history of stroke (105,106). Several lines of evidence also revealed that the level of GDF15 was found to be positively associated with the severity of ischemic stroke, such that GDF15 can be used as a biomarker for predicting an unfavorable outcome 90 days after the stroke event (107,108). Plasma GDF15 concentration on admission has been reported to serve as an independent prognostic biomarker of mortality in patients with ischemic stroke following acute revascularization therapy (17). A previous study also investigated the relationship between GDF15 and in 254 patients with hypertension who suffered from stroke for the first, which found that GDF15 can be used as an independent predictor of stroke in individuals without any prior history of stroke (109). In addition, it was reported that GDF15 mRNA expression was markedly

upregulated in the hippocampus and parietal cortex of mice at 3 and 24 h after middle cerebral artery occlusion (110). This suggests that GDF15 can participate in the regulation of post-lesion responses, further supporting the notion that GDF15 participates in the occurrence and development of cerebral stroke (110).

**PD.** The main property of PD is the degeneration and loss of dopaminergic neurons in the substantia nigra and nigrostriatal pathway, which is caused by the formation of Lewy bodies as a result of aberrant  $\alpha$ -synuclein deposition (111). PD is characterized by symptoms of dyskinesia, including tremor, stiffness, slow motion and unstable posture (111). PD also has a complex and multifactorial pathological process, which typically involves the aggregation of  $\alpha$ -synuclein, oxidative stress, mitochondrial dysfunction, iron deposition and neuronal apoptosis (112). Maetzler *et al* (113) previously revealed that GDF15 exhibited a positive correlation with the age of PD symptom onset, Hoehn and Yahr scale score and expression of the neurodegenerative marker Tau. GDF15 was also identified to be an independent risk factor for Unified Parkinson's Disease Rating Scale-III score through multiple linear regression analysis, where subsequent receiver operating characteristic curve analysis revealed that GDF15 exhibited a sensitivity of 71.15% and a specificity of 87.50% for the detection of PD (114,115). However, since GDF15 levels exhibit substantial overlap between patients with PD and healthy individuals, this marker alone may not be sufficient as a diagnostic tool. However, these aforementioned findings indicate collectively that GDF15 expression is closely associated with PD.

The neurotoxin 6-hydroxydopamine (6-OHDA) can be taken up preferentially by dopaminergic and noradrenergic transporters, leading to the degeneration of catecholaminergic neurons (116). Strelau *et al* (84) demonstrated that unilateral injections of GDF15 into the medial forebrain bundle immediately above the substantia nigra prior to 6-OHDA

administration were able to confer protection against complete lesion formation induced by 6-OHDA, which prevented the loss of the dopaminergic neurons. Consistently, Machado *et al* (117,118) further demonstrated that endogenous GDF15 may promote the survival of dopaminergic neurons by regulating the inflammatory response after 6-OHDA-induced brain injury. GDF15 released by astrocytes exerted a protective effect on vulnerable nigral neurons during PD and on induced pluripotent stem cell-derived dopaminergic neurons subjected to 1-methyl-4-phenylpyridinium toxicity, which may explain the selective degeneration or protection of dopaminergic neurons in PD because GDF15 is expressed 230-fold higher in the neighboring ventral tegmental area astrocytes than the substantia nigra pars compacta (20,119). Mitochondrial dysfunction is another possible mechanism that has been proposed to serve a role in PD (112). The HT22 hippocampal cell line is considered to be a suitable model for studying PD. GDF15 overexpression was shown to reverse the effects of oxygen consumption, cell viability and mitochondrial membrane potential caused by oligomycin in HT22 cells, where further study revealed that GDF15 may regulate mitochondrial membrane potential and oxygen consumption through the PI3K/AKT signaling pathway (120). Furthermore, GDF15 was reported to be a more sensitive measure for diagnosing mitochondrial dysfunction compared with that of lactate stress test in Japanese patients with PD (121). Collectively, these findings suggest that GDF15 may promote the survival of dopaminergic neurons and exerts a protective effect by preserving normal mitochondrial function.

## 6. Conclusion and future directions

GDF15 is widely expressed in brain tissues and has been found to be involved in the pathophysiological processes underlying a number of brain disorders, particularly in AD, cerebral stroke and PD. Although progress has been made over the past decade, several unresolved problems remain. The specific signaling pathways mediated by GDF15 in AD have not been fully elucidated. In addition, the reference range and sensitivity of GDF15 as a biomarker for the prognosis and risk stratification of related diseases have yet to be determined. Furthermore, it remains unknown if there are other GDF15 receptors involved in mediating the pathophysiological processes of brain disorders in addition to the GFRAL receptor. Whether the plasma concentration of GDF15 can be artificially regulated to treat brain disorders is also a question that requires further study. Therefore, addressing these questions aforementioned may provide further clues for the prevention and treatment of these brain disorders.

## Acknowledgements

Not applicable.

## Funding

The authors acknowledge the financial supports from the Natural Science Foundation of Hunan Province, (grant nos. 2019JJ40249 and 2018JJ3455), Outstanding Young Aid Program for Education Department of Human Province (grant no. 18B274), the Major Project of Social Science

Achievement Review Committee in Hunan Province (grant no. XSP20ZDI013), Key Project of Hunan Provincial Department of Education (grant no. 20A427).

## Availability of data and materials

Not applicable.

## Authors' contributions

WWJ, ZZZ, PPH, XPO contributed to data gathering, manuscript drafting, critical revision of the manuscript. LPJ, JZC, XTZ, MH and YKZ revised the manuscript. All authors read and approved the final manuscript. Data authentication is not applicable.

## Ethics approval and consent to participate

Not applicable.

## Patient consent for publication

Not applicable.

## Competing interests

The authors declare that they have no competing interests.

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